Hauda, K.

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46

(2) INFORMATION FOR SEQ ID NO:1:

1632

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/982,284

This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING		
2 3 4	(1) G	General Information: ENTE	R	
5 6 7 8 9	(i)	APPLICANT: LUBON, Henryk PALEYANDA, Rekha DROHAN, William VELANDER, William		
10 11 12 13	(ii)	TITLE OF INVENTION: METHODS FOR THE DEGRADATION AND DETOXIFICATION OF ORGANIC MATERIAL USING URINE PRODUCED TRANSGENIC ANIMALS AND RELATED TRANSGENIC ANIMALS AND PROTEINS		- ‡
14 15 16	(iii)	NUMBER OF SEQUENCES: 3	00 APR	ECHOS
17 18 19 20 21 22 23	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: FOLEY & LARDNER (B) STREET: 3000 K Street, N.W. (C) CITY: Washington (D) STATE: D.C. (E) COUNTRY: U.S.A. (F) ZIP: 20007-5109	PR -6 AN 9:35	
25 26 27 28 29	(v)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30		80
30 31 32 33 34 35	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: US 08/982,284 (B) FILING DATE: 01-DEC-1997 (C) CLASSIFICATION:		
36 37 38 39	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: Granados, Patricia D. (B) REGISTRATION NUMBER: 33,683 (C) REFERENCE/DOCKET NUMBER: 030523/0141		
40 41 42 43 44 45	(ix)	TELECOMMUNICATION INFORMATION: (A) TELEPHONE: (202) 672-5300 (B) TELEFAX: (202) 672-5399		

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/982,284

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47		
48	(i) SEQUENCE CHARACTERISTICS:	
49	(A) LENGTH: 69 base pairs	
50	(B) TYPE: nucleic acid	
51	(C) STRANDEDNESS: single	
52	(D) TOPOLOGY: linear	
53	(1)	
54	(ii) MOLECULE TYPE: DNA (genomic)	
55	(11, 1,0110011 11111 11111 (3010)	
56		
57		
58		
59	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
60	(XI) BEQUENCE DESCRIPTION: BEQ ID NO.I:	
61	GGTACGTCGA CACAGGTGCC AGTGCCTCCA GAATGTGGCA GCTCACAAGC CTCCTGCTGT	60
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63	TCCTTCCCCC	69
	TCGTGGCCA	69
64	(2) INFORMATION FOR GEO. ID NO. 2.	
65	(2) INFORMATION FOR SEQ ID NO:2:	
66	(+) GEOURNGE GUADAGMEDIGMIGG	
67	(i) SEQUENCE CHARACTERISTICS:	
68	(A) LENGTH: 22 base pairs	1.1
69	(B) TYPE: nucleic acid	9 9
70	(C) STRANDEDNESS: single	- C
71	(D) TOPOLOGY: linear	9 177
72		OD APR-6
73	(ii) MOLECULE TYPE: other nucleic acid	0
74	(A) DESCRIPTION: /desc = "Primer"	الورد أ
75		三 具 四
76		
77		ر م
78		-6 图 9: 35
79	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	J . (
80		
81	CCCAGGCTCA GCGCACTCAT CC	22
82		
83	(2) INFORMATION FOR SEQ ID NO:3:	
84		
85	(i) SEQUENCE CHARACTERISTICS:	
86	(A) LENGTH: 21 base pairs	
87	(B) TYPE: nucleic acid	
88	(C) STRANDEDNESS: single	
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91	(ii) MOLECULE TYPE: other nucleic acid	
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93		
94		
95		
96		
97	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
98		
99	GTCACAGCAA TGCCACCTGA C	21

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SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/982,284

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